



Preface

Epigenetics

Epigenetics as coined by the British biologist Conrad Waddington to explain differentiation [1] deals with alterations in gene expression unrelated to changes in DNA sequences. Although not based on DNA mutations, epigenetic changes are clonally inheritable from one cell to the other, potentially modified by environmental stimuli and even transmitted from parent to offspring in some instances. Epigenetic processes play an important role in the earliest stages of development when epigenetic modifications of chromatin structure, such as DNA methylation, cause lifelong and sometimes trans-generational traits [2]. Besides DNA methylation, specific epigenetic mechanisms among others comprise histone modifications (e.g. by Polycomb–Trithorax protein complexes), ATP-dependent chromatin remodeling and non-coding RNA-mediated gene-silencing. Most of these epigenetic processes have an impact on chromatin organization and maintenance, and occur in many species affecting important biological processes such as genomic imprinting, X-inactivation, reprogramming in embryogenesis, and chromosomal structures. The past few years have seen an explosion of knowledge in the molecular mechanisms of epigenetics, along with the realization that epigenetic processes could also result in phenotypic variation [3]. Given that diseases are phenotypes, it is not surprising that impaired epigenetic processes are a major fundamental risk in acquired disorders such as cancer and thus epigenetic drug-based chemotherapeutic approaches are subjects of intense research [4].

The present Special Issue on EPIGENETICS comprises 16 reviews covering the topics of *DNA methylation* [5–7], *histone modification and chromatin regulation* [8,9], *environmental epigenetics* [10–12], *developmental plasticity* [13,14], *cancer epigenetics* [15–18] and *epigenetics and therapeutic potential* [19,20]. The editors thank the authors and reviewers for their efforts and hope that the readers will enjoy this updated, comprehensive and foresighted discussion on ongoing epigenetic research.

References

- [1] Waddington, C.H. (1942) The epigenotype. *Endeavour* 1, 18–20.
- [2] Skinner, M.K. (2011) Environmental epigenetic transgenerational inheritance and somatic epigenetic mitotic stability. *Epigenetics* 6, 838–842.
- [3] Whitelaw, E. and Martin, D.I. (2001) Retrotransposons as epigenetic mediators of phenotypic variation in mammals. *Nat. Genet.* 27, 361–365.
- [4] Taby, R. and Issa, J.-P.J. (2010) Cancer epigenetics. *CA Cancer J. Clin.* 60, 376–392.
- [5] Meyer, P. (2011) DNA methylation systems and targets in plants. *FEBS Lett.* 585, 2008–2015.
- [6] Pelizzola, M. and Ecker, J.R. (2011) The DNA methylome. *FEBS Lett.* 585, 1994–2000.
- [7] Scharf, A.N.D. and Imhof, A. (2011) Every methyl counts – Epigenetic calculus. *FEBS Lett.* 585, 2001–2007.
- [8] Atanassov, B.S., Koutelou, E. and Dent, S.Y. (2011) The role of deubiquitinating enzymes in chromatin regulation. *FEBS Lett.* 585, 2016–2023.
- [9] Di Lorenzo, A. and Bedford, M.T. (2011) Histone arginine methylation. *FEBS Lett.* 585, 2024–2031.
- [10] Caldj, C., Hellstrom, I.C., Zhang, T.-Y., Diorio, J. and Meaney, M.J. (2011) Environmental regulation of the neural epigenome. *FEBS Lett.* 585, 2049–2058.
- [11] Feser, J. and Tyler, J. (2011) Chromatin structure as a mediator of aging. *FEBS Lett.* 585, 2041–2048.
- [12] Turner, B.M. (2010) Environmental sensing by chromatin: An epigenetic contribution to evolutionary change. *FEBS Lett.* 585, 2032–2040.
- [13] Radford, E.J., Ferrón, S.R. and Ferguson-Smith, A.C. (2011) Genomic imprinting as an adaptive model of developmental plasticity. *FEBS Lett.* 585, 2059–2066.
- [14] Prezioso, C. and Orlando, V. (2011) Polycomb proteins in mammalian cell differentiation and plasticity. *FEBS Lett.* 585, 2067–2077.
- [15] Cheung, N. and So, C.W.E. (2011) Transcriptional and epigenetic networks in haematological malignancy. *FEBS Lett.* 585, 2100–2111.
- [16] Estéio, M.R. and Issa, J.-P.J. (2011) Dissecting DNA hypermethylation in cancer. *FEBS Lett.* 585, 2078–2086.
- [17] Maradeo, M.E. and Cairns, P. (2011) Translational application of epigenetic alterations: Ovarian cancer as a model. *FEBS Lett.* 585, 2112–2120.
- [18] Melo, S.A. and Esteller, M. (2011) Dysregulation of microRNAs in cancer: Playing with fire. *FEBS Lett.* 585, 2087–2099.
- [19] Huang, Y.-W., Kuo, C.-T., Stoner, K., Huang, T.H.-Y. and Wang, L.S. (2011) An overview of epigenetics and chemoprevention. *FEBS Lett.* 585, 2129–2136.
- [20] Sugii, S. and Evans, R.M. (2011) Epigenetic codes of PPAR γ in metabolic disease. *FEBS Lett.* 585, 2121–2128.

Jean-Pierre Issa

Department of Leukemia, The University of Texas,
MD Anderson Cancer Center, 1515 Holcombe Boulevard,
Houston, TX 77030, USA

E-mail address: jissa@mdanderson.org

Wilhelm Just

Biochemie-Zentrum der Universität Heidelberg (BZH),
Im Neuenheimer Feld 328, D-69120 Heidelberg,
Germany

E-mail addresses: wilhelm.just@bzh.uni-heidelberg.de

Available online 17 June 2011